BGL4 Beta-Glucosidase and Nucleic Acids
Encoding the Same
Dunn-Coleman et al.
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Figure 1-A T. reesei bgl4 nucleotide sequence (non-coding is in bold, coding is in regular font)

ACAAGTTCAAGATCGAGTTCGGCTCCGCACCCACCTACACCCTCAAGGGCGACACCATCGTCCCGGCCACGGCTCCTCC GCGTCGGCGGCTGCAAGGTCATTGACGACCAGGCCGAAATCGAAAAGTCCGTCGCCCTCGCCAAGGAGCACGACCAGGTCA AGCAGGTCCCCGGCGATGCCTTCTTCGGCTCCGCCACCCGCGAGGAGGGGCCGCCGCATCAATCTCGTCAAGGGCAACACGT TCCACGTCATTGACCAGAGGGCTAGGGAAGTTCTTCAGTTCGTCAAGAAGTGTGCTGCCTCCGGAGTGAGGAGAACGGCC ACGAGAACAACGTTCTGCCCTTGAGCAAGAAGAAGACGCTGATTGTCGGCCCCCAACGCCAAGGAGGCCAATACCACG GCTGGAGGGTCTTCAACGAGCCCCCTGGTACCCCTAACCGCCAGCACATTGACGAGCTCTTCTTCACCAAGACGGACATGC ACCTGGTGGACTACTACCACCCCAAGGCGGCAGACACGTGGTACGCCGACATGGAGGGCACGTACACCGCCGACGAGGGACT TTCGAAAGGAATGGGGTTGGGATGGCCTAATCATGAGCGACTGGTACGGCACATACAGTACCACAGAAGCCGTTGTGGCCAG GCCTCGACCTCGAGATGCCCGGGACCTCCCACGCTTCCGAGGAGAACACTCCAAGTTCAACGTCTCCAACGGAAAGCCCTTTA CCGAGACGACTGTCAACACCCCCCGGAAACGGCAGCTCTCCTCCGGAAGGTTGGCAACGAGGGCATCGTGCTGCTGAAGA CGIACACCGICGGCGCCIACACCACCGTICCICCCAIICIAGGCGAGCAGIGCCICACGCCCGACGGCGCTCCGGGCAIGC CGCCATATAGAGCAGCATATTCTTCTATCTCCAAAGATCCCCTCACCGAGAGTTCTATTCACCCGACCCTTGCCTTGTCAT **CCAGTCCTTCCATC**ATGGCTGATATTGATGTTGAGGCCCATCTTGAAGAAGCTCACCCTGGCCGAGAAGGTCGATCTGCTGG CTGGTATCGACTTCTGGCACACAAAGGCTCTCCCCAAGCATGGAGTCCCCTCTCTCCGCTTTACAGATGGCCCCAACGGCG ACATGCAACGCTCCCCTCTCGGTGGACGTGGCTTCGAGTCGATTGGTGAGGATCCGTTCCTGGCGGGCTTGGGAGCTGCGG TGATGGTGCAGAGCATCGTCACGGAGCGGGCTCTCCGTGAAATCTACGCACTCCCGTTCCAGATTGCTGTGCGAGACTCCC AGCCGGGTGCGTTCATGACGGCGTACAATGGCATCAATGGCGTGTCGTGCAGCGAGAACCCTAAATATCTTGATGGGATGC TAAGAGGGACCAAGTTCTTCAATGGCGTCCCTGCGGCCTGCTTCCCTTGCGGCACGTCGCTCGGTTCCACATTCAACCAAA CICTCATCCGCGCGCATTCAGAGCACTGGAGTGCAGGCTACGATCAAGCACTTTTTGTGCAATGATCAGGAGGACAGGCGCG TTATAGTCGCTTTGTTAAATTGGCCTCGAGGTCGACCCACGCGTCCGGCTGCTTGTCCCGTTCCTGTGCCTGATGTCTATC CCCCCTTCGTTGCTCTGGTGTGTCCAAGGTCTACCCTGCAGTGGTTTTTGAACCCTTGATATCCTGCTTGAGCATCCGCGT CTCTGCTCGAAGAGGCAGGTAAGATGATGGGCCAAAGAGGCCCATCGCTAAGAGTGCGCATGTGATCCTCGGCCCGACTATCA

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TGAGCGACAGCAGCGCAGCGAAGGATGGGGTTGCGCTCAGGGGTAAGTTTACGGTGGGAGAGACGTATTGGTGGTCTGGCG CCCTCTCCGTGAAGAACACCGGGTTCCGTGCCCGGCGCACAGGTGGCCCCAGCTCTACGTCAAGCCCCTCCAAGGAGC TTAACCGCCCCGTCAAGGAGCTCAAGGGCTTCGCAAAGGTCGAACTGCAGCCCGGCGAGACGAAGGCGGTGACAATCGAGG AGCAGGAGAAGTACGTCGCTGCGTATTTTGATGAGGAGCGGGATCAGTGGTGTGTCGAAAAGGGGTGACTATGAGGTTATCG TGTAAAGTCGTGCATCATCTTTGGCAGATTGAATCCAGTCACTTTGCATATAGCGCCGATGAAATGAGAACCAACGATCAT TCGACGCCACGCCCGCCGTCATCCAGGCCTGGTACGGCGCCAACGAGACGGGCAACTCCATTGCCGACGTCGTCTTTGGCG ACTACAACCCCTCGGGCAAGCTGTCCCTCAGCTTCCCCAAGCGCCTGCAGGACAACCCCGCGTTTCTCAACTTCCGCACCG AGGCCGGGCGCACGCTGTACGGCGAGGACGTCTACGTCGGGTACAGGTACTACGAGTTTGCCGACAAGGACGTCAATTTCC CCTTTGGCCACGGCCTGTCCTACACCACTTTTGCCTTTTCCAATCTCTCCGTGTCTCACAAGGACGGCAAGCTGAGCGTGTGT AGCTCATTGCCGACGTGGCCGCCGCGAACCCCAAACACGTCGTCGTCATGCAGACGGGCACCCCCGAGGAGATGCCTGGC BGL4 Beta-Glucosidase and Nucleic Acids Encoding the Same Dunn-Coleman et al. SN# Unassigned Docket No. GC696 Sheet 3 of 4

Figure 2-A Predicted T. reesei BGL4 amino acid sequence

MADIDVEAIL KKLTLAEKVD LLAGIDFWHT KALPKHGVPS LRFTDGPNGV RGTKFFNGVP AACFPCGTSL	GSTFNQTLLE EAGKMMGKEA IAKSAHVILG PTINMQRSPL GGRGFESIGE DPFLAGLGAA ALIRGIQSTG	VQATIKHFLC NDQEDRRMMV QSIVTERALR EIYALPFQIA VRDSQPGAFM TAYNGINGVS CSENPKYLDG	MLRKEWGWDG LIMSDWYGTY STTEAVVAGL DLEMPGPPRF RGETLKFNVS NGKPFIHVID QRAREVLQFV	KKCAASGVTE NGPETTVNNT PETAALLRKV GNEGIVLLKN ENNVLPLSKK KKTLIVGPNA KQATYHGGGS	AALRAYYAVT PFDGLSKQLE TPPSYTVGAY TTVPPILGEQ CLTPDGAPGM RWRVFNEPPG TPNRQHIDEL	FFTKTDMHLV DYYHPKAADT WYADMEGTYT ADEDCTYELG LVVCGTAKAY VDDQLVVDNA TKQVPGDAFF	GSATREETGR INLVKGNTYK FKIEFGSAPT YTLKGDTIVP GHGSLRVGGC KVIDDQAEIE KSVALAKEHD	QVIICAGLNA DWETEGADRA SMKLPGVLDQ LIADVAAANP NTVVVMQTGT PEEMPWLDAT PAVIQAWYGG
10 20 30 40 50	80 90 100 110 120	150 160 170 180 200	220 230 240 250 280	350 330 340 350	360 370 380 390 400	430 440 450 460 470 480	500 510 520 530 540	570 580 590 630

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Predicted T. reesei BGL4 amino acid sequence Figure 2-B

700 700	2PG 770	
EF ADKDVNFFF	LK GFAKVELQP(WW SGV
690 70	760	830 833
YGED VYVGYRYY	LQAA KINRPVKE	ALRG KFTVGETY
680	750	820
NETGNSIADV VFGDYNPSGK LSLSFPKRLQ DNPAFLNFRT EAGRTLYGED VYVGYRYYEF ADKDVNFFFG	HGLSYTTFAF SNLSVSHKDG KLSVSLSVKN TGSVPGAQVA QLYVKPLQAA KINRPVKELK GFAKVELQPG	ETKAVTIEEQ EKYVAAYFDE ERDQWCVEKG DYEVIVSDSS AAKDGVALRG KFTVGETYWW SGV
640 650 650 700	710 720 730 740	780 830 83
LSFPKRLQ DNPA	SVSLSVKN TGSV	DQWCVEKG DYEV
660	730	800
FGDYNPSGK LSI	NLSVSHKDG KL.	KYVAAYFDE ER
650	720	790
NETGNSIADV V	HGLSYTTFAF S 710	ETKAVTIEEQ F

790